

Figure 1

Sequence alignment of mouse Serca 1, 2 and 3 protein.

5	Sercala	1	MEAAHSKSTEECLSYFGVSETTGLTPDQVKRHL	EKYGPNELPAEEGKSLWELVVEQFEDL
	Serca2a	1	..N..T.TV..V.GH..N.S..SLE..	KLK.RW.S.....T.L...I.....
	Serca2b	1	..N..T.TV..V.GH..N.S..SLE..	KLK.RW.S.....T.L...I.....
	Serca3a	1	..E..LL.AADV.RR.S.TAEG..SLE..	TDAR.R.....T.....
	Serca3b	1	..E..LL.AADV.RR.S.TAEG..SLE..	TDAR.R.....T.....
10	Serca3c	1	..E..LL.AADV.RR.S.TAEG..SLE..	TDAR.R.....T.....
	Sercala	61	LVRILLAA	CISFVLAWFEEGEETVTAFVEPFVILLILIANAIVGVWQERNAENAIEALK
	Serca2a	61	I.....V.....
	Serca2b	61	I.....V.....
15	Serca3a	61LV	T.....L.M..V.....S.....
	Serca3b	61LV	T.....L.M..V.....S.....
	Serca3c	61LV	T.....L.M..V.....S.....
	Sercala	121	EYEPEMGK	VYRADRKSVQRIKARDIVPGDIVEAVGDKVPADIRILSIKSTTLRVDQSIL
20	Serca2a	121Q.....K.....I.....LT.....
	Serca2b	121Q.....K.....I.....LT.....
	Serca3a	121I.S..G..R.....L.LIE.....
	Serca3b	121I.S..G..R.....L.LIE.....
	Serca3c	121I.S..G..R.....L.LIE.....
25	Sercala	181	TGESVSVIKHTDPVPDPRAVNQDKKNMLFSGTNIAAGKAVGIVATTGVSTEIGKIRDQMA	
	Serca2a	181	M.V.VA..N.....E.V
	Serca2b	181	M.V.VA..N.....E.V
	Serca3a	181T...AIS..L.VAVA..LQ..L...S...
30	Serca3b	181T...AIS..L.VAVA..LQ..L...S...
	Serca3c	181T...AIS..L.VAVA..LQ..L...S...
	Sercala	241	ATEQDKTPLQQKLDEFGEQLSKVISLICVAVWLINIGHFNDPVHGGSWFRGAIYYFKIAV	
	Serca2a	241ER.....I..I.....I.....
35	Serca2b	241ER.....I..I.....I.....
	Serca3a	241	..V.PER....R..HA..V....V.....A..A.....L..V	
	Serca3b	241	..V.PER....R..HA..V....V.....A..A.....L..V	
	Serca3c	241	..V.PER....R..HA..V....V.....A..A.....L..V	
40	Sercala	301	ALAVAAIPEGLP	AVITCLALGTRRMAKKNAIVRSLPSVETLGCTSVICSDKTGTLTTNQ
	Serca2a	301	
	Serca2b	301	
	Serca3a	301	R.....
	Serca3b	301	R.....
45	Serca3c	301	R.....
	Sercala	361	MSVCKMFIIDKVDGDVCSLNEFSITG	STYAPEGEVLKNDKPVRAGQYDGLVELATICALC
	Serca2a	361R..L..E..T.....	I..Q.D...KCH.....
	Serca2b	361R..L..E..T.....	I..Q.D...KCH.....
50	Serca3a	361R..VVAEAEAGT.R.H..T.S.T..T.....	RQGEQ..C..F.....
	Serca3b	361R..VVAEAEAGT.R.H..T.S.T..T.....	RQGEQ..C..F.....
	Serca3c	361R..VVAEAEAGT.R.H..T.S.T..T.....	RQGEQ..C..F.....
	Sercala	421	NDSSLDFNETKG	VYEKVGEATETALTLVEKMNVFNT
55	Serca2a	421	...A..Y..A.....	LSKVERANACNSVIQLMK
	Serca2b	421	...A..Y..A.....	C.....D..LKG..I.....K...
	Serca3a	421	...A..Y..A.....	C.....D..LKG..I.....K...
	Serca3b	421	...A..Y..A.....	C.....D..LKG..R.....G.....K...R
	Serca3c	421	...A..Y..A.....	C.....D..LKG..R.....G.....K...R

	Sercala	481	KEFTLEFSRDRKMSVYCPAKSSRAAVGNKMFVKGAPEGVIDRCNYVRVGTTRVPLTGP	
	Serca2a	481T.N.P..TSMS-.....	THI..S.K..M.PG
	Serca2b	481T.N.P..TSMS-.....	THI..S.K..M.PG
5	Serca3a	481T.TRADPKVQ.S.....	S..E..SS...SRTA..STT
	Serca3b	481T.TRADPKVQ.S.....	S..E..SS...SRTA..STT
	Serca3c	481T.TRADPKVQ.S.....	S..E..SS...SRTA..STT
	Sercala	541	VKEKIMSVIKEWGTGRDTLRLALATRDTPPKREEMVLDDSAKFMEYEMDLTFVGVVGML	
	Serca2a	540	..Q.....R...S.S.....H.N.L....H.E...N.IK..TN.....C....	
10	Serca2b	540	..Q.....R...S.S.....H.N.L....H.E...N.IK..TN.....C....	
	Serca3a	541	SR.H.LAK.RD..S.S.....RK.D.H...CSR.VQ..T.....C....	
	Serca3b	541	SR.H.LAK.RD..S.S.....RK.D.H...CSR.VQ..T.....C....	
	Serca3c	541	SR.H.LAK.RD..S.S.....RK.D.H...CSR.VQ..T.....C....	
15	Sercala	601	DPPRKEVTGSIQLCRDAGIRVIMITGDNKGTAIAICRRIGIFSENEEVTDRAYTGREFDD	
	Serca2a	600I..AS.VK...Q.....V.....GQD.D..SK.F.....E	
	Serca2b	600I..AS.VK...Q.....V.....GQD.D..SK.F.....E	
	Serca3a	601P..AAC.TR.SR....V.....V....L...GDT.D.LGK.....	
	Serca3b	601P..AAC.TR.SR....V.....V....L...GDT.D.LGK.....	
20	Serca3c	601P..AAC.TR.SR....V.....V....L...GDT.D.LGK.....	
	Sercala	661	LPLAEQREACRACC FARVEPSHKS KIVEYLQSYDEITAMTGDGVNDAPALKKAEIGIAM	
	Serca2a	660	.SPSA..D..LN.R.....F...F.....S.....	
	Serca2b	660	.SPSA..D..LN.R.....F...F.....S.....	
25	Serca3a	661	.SPEQ..Q...T.R.....A..R..N..FN.....	
	Serca3b	661	.SPEQ..Q...T.R.....A..R..N..FN.....	
	Serca3c	661	.SPEQ..Q...T.R.....A..R..N..FN.....	
	Sercala	721	GSGTAVAKTASEMVLADDNFSTIVAAVEEGRAIYNNMKQFIRYLISSNVGEVVCIFLTAA	
30	Serca2a	720	
	Serca2b	720	
	Serca3a	721S.A....S....AS.....I	
	Serca3b	721S.A....S....AS.....I	
	Serca3c	721S.A....S....AS.....I	
35	Sercala	781	LGLPEALIPVQLLWVNLT DGLPATALGFNPPDLDIMDRPPRSPKEPLISGWLFFRYMAI	
	Serca2a	780	..F.....NK..N.....L..	
	Serca2b	780	..F.....NK..N.....L..	
	Serca3a	781EK..N.R.A.....L..	
40	Serca3b	781EK..N.R.A.....L..	
	Serca3c	781EK..N.R.A.....L..	
	Sercala	841	GGYVGAATVGAAAWFLYAE DGP HVSYHQLTHFMQCTEHNPEFDGLDCEVFEA PEPMTMA	
	Serca2a	840	.C.....IA.DG..R..FY..S..L..K.D..D..V..AI..S.Y.....	
45	Serca2b	840	.C.....IA.DG..R..FY..S..L..K.D..D..V..AI..S.Y.....	
	Serca3a	841	.V...L...A..T.....DAE..Q.TFY..RN.LK.S.D..L.A.I..K..SRF.T..	
	Serca3b	841	.V...L...A..T.....DAE..Q.TFY..RN.LK.S.D..L.A.I..K..SRF.T..	
	Serca3c	841	.V...L...A..T.....DAE..Q.TFY..RN.LK.S.D..L.A.I..K..SRF.T..	
50	Sercala	901	LSVLVTIEMCNALNSLSENQSLLRMPPWVNWI LLSICLMSLHFLILYVDPLPMIFKLR	
	Serca2a	900E....V.....E...L..QIT	
	Serca2b	900E....V.....E...L..QIT	
	Serca3a	901V.....L.P....AVVM..A.....L.P...L..QVT	
	Serca3b	901V.....L.P....AVVM..A.....L.P...L..QVT	
55	Serca3c	901V.....L.P....AVVM..A.....L.P...L..QVT	

Sercala 961 ALDFTQWLMVLKISLPVIGLDELLKFIARNYLEG
Serca2a 960 P.NL.....LM..T...V.....QPAILE
Serca2b 960 P.NL.....LM..T...V.....QPGKECVQPATKSSCSLSACTDGISWP
Serca3a 961 P.SGR..GV..QM....L...A..YLS..HMDEKKDLK
5 Serca3b 961 P.SGR..GV..QM....L...A..YLS..HMD.VLGTFMQARSQQLPTTSRTPYHTGKK
Serca3c 961 P.SGR..GV..QM....L...A..YLS..HMD.VLGTFMQARSQQLPTTSRTPYHTGLA

10 Serca2b 1020 FVLLIMPLVVWVYSTDTNFSDMFWS
Serca3b 1021 GPEVNPGSRGESPVWPSD
Serca3c 1021 SWKKRT

Figure 2

Sequence similarity of Serca2 proteins in mammalian species

	Mouse_2a	1	MENAHTKTVEEVLGHFGVNESTGLSLEQVKKLKERWGSNELPAEEGKTLLELVIEQFEDL
5	Mouse_2b	1
	Rat_2b	1
	Rat_2a	1
	Dog_2a	1
	Cat_2a	1Y.....
10	Pig_2a	1
	Pig_2b	1
	Human_2b	1
	Human_2c	1
	Human_2a	1
15	Rabbit_2a	1
	Rabbit_2b	1
	Mouse_2a	61	LVRILLLAACISFVLAWFEEGEETITAFVEPFVILLILVANAIIVGVWQERNAENATEALK
	Mouse_2b	61
20	Rat_2b	61
	Rat_2a	61
	Dog_2a	61
	Cat_2a	61
	Pig_2a	61
25	Pig_2b	61
	Human_2b	61
	Human_2c	61
	Human_2a	61
	Rabbit_2a	61
30	Rabbit_2b	61
	Mouse_2a	121	EYEPEMGKVYRQDRKSVQRIKAKDIVPGDIVEIAVGDKVPADIRLTSIKSTTLRVDQSIL
	Mouse_2b	121
	Rat_2b	121
35	Rat_2a	121
	Dog_2a	121
	Cat_2a	121
	Pig_2a	121
	Pig_2b	121
40	Human_2b	121
	Human_2c	121
	Human_2a	121
	Rabbit_2a	121
	Rabbit_2b	121
45	Mouse_2a	181	TGESVSVIKHTDPVPDPRAVNQDKKNMLFSGTNIAAGKAMGVVVATGVNTEIGKIRDEMV
	Mouse_2b	181
	Rat_2b	181
	Rat_2a	181
50	Dog_2a	181
	Cat_2a	181
	Pig_2a	181
	Pig_2b	181
	Human_2b	181
55	Human_2c	181
	Human_2a	181
	Rabbit_2a	181
	Rabbit_2b	181

	Mouse_2a	241	ATEQERTPLQQKLDEFGEQLSKVISLICIAVWIINIGHFNDPVHGGSWIRGAIYYFKIAV
	Mouse_2b	241
5	Rat_2b	241
	Rat_2a	241
	Dog_2a	241
	Cat_2a	241
	Pig_2a	241
	Pig_2b	241
10	Human_2b	241
	Human_2c	241
	Human_2a	241
	Rabbit_2a	241
	Rabbit_2b	241
15	Mouse_2a	301	ALAVAAIPEGLPAVITCLALGTRRMAKKNAIVRSLPSVETLGCTSVICSDKTGTLTTNQ
	Mouse_2b	301
	Rat_2b	301
	Rat_2a	301
20	Dog_2a	301
	Cat_2a	301
	Pig_2a	301
	Pig_2b	301
	Human_2b	301
25	Human_2c	301
	Human_2a	301
	Rabbit_2a	301
	Rabbit_2b	301
30	Mouse_2a	361	MSVCRMFIIDKVEGDTCSLNEFSITGSTYAPIGEVQKDDKPVKCHQYDGLVELATICALC
	Mouse_2b	361
	Rat_2b	361 T
	Rat_2a	361 T
	Dog_2a	361 R .. S .. T .. H ..
35	Cat_2a	361 T .. H ..
	Pig_2a	361 T .. H ..
	Pig_2b	361 T .. H ..
	Human_2b	361 R .. T .. H .. N ..
	Human_2c	361 R .. T .. H .. N ..
40	Human_2a	361 R .. T .. H .. N ..
	Rabbit_2a	361 D .. T .. H ..
	Rabbit_2b	361 D .. T .. H ..
	Mouse_2a	421	NDSALDYNEAKGVYEVGEATETALTCLVEKMNVFDTELKGLSKIERANACNSVIKQLMK
45	Mouse_2b	421
	Rat_2b	421
	Rat_2a	421
	Dog_2a	421
	Cat_2a	421 K. F ..
50	Pig_2a	421
	Pig_2b	421
	Human_2b	421
	Human_2c	421
	Human_2a	421
55	Rabbit_2a	421
	Rabbit_2b	421

	Mouse_2a	481	KEFTLEFSRDRKSMHSVYCTPNKPSRTSMSKMFVKGAPEGVIDRCTHIRVGSTKVPMTPGV
	Mouse_2b	481
	Rat_2b	481
	Rat_2a	481
5	Dog_2a	481
	Cat_2a	481
	Pig_2a	481
	Pig_2b	481
	Human_2b	481
10	Human_2c	481
	Human_2a	481
	Rabbit_2a	481
	Rabbit_2b	481
15	Mouse_2a	541	KQKIMSVIREWGSGSDTLRCLALATHDNPLKREEMHLEDSANFIKYETNLTFVGCVGMLD
	Mouse_2b	541
	Rat_2b	541
	Rat_2a	541
	Dog_2a	541V.....
20	Cat_2a	541V.....
	Pig_2a	541
	Pig_2b	541
	Human_2b	541
	Human_2c	541
25	Human_2a	541
	Rabbit_2a	541
	Rabbit_2b	541
	Mouse_2a	601	PPRIEVASSVLCRQAGIRVIMITGDNKGTAVAICRRIGIFGQDEDVTSKAFTGREFDEL
30	Mouse_2b	601
	Rat_2b	601
	Rat_2a	601
	Dog_2a	601
	Cat_2a	601
35	Pig_2a	601
	Pig_2b	601
	Human_2b	601
	Human_2c	601
	Human_2a	601
40	Rabbit_2a	601
	Rabbit_2b	601
	Mouse_2a	661	SPSAQRDACLNRCAFARVEPSHKSKIVEFLQSFDEITAMTDGVNDAPALKSEIGIAMG
	Mouse_2b	661
45	Rat_2b	661
	Rat_2a	661
	Dog_2a	661
	Cat_2a	661
	Pig_2a	661	N.....E.....
50	Pig_2b	661	N.....E.....
	Human_2b	661	N.....
	Human_2c	661	N.....
	Human_2a	661	N.....
	Rabbit_2a	661	N.....
55	Rabbit_2b	661	N.....

Mouse_2a	721	SGTAVAKTASEMVLADDNFSTIVAAVEGRAIYNNMKQFIRYLISSNVGEVVCIFLTAAL
Mouse_2b	721
Rat_2b	721
Rat_2a	721
5 Dog_2a	721
Cat_2a	721
Pig_2a	721
Pig_2b	721
Human_2b	721
10 Human_2c	721
Human_2a	721
Rabbit_2a	721
Rabbit_2b	721
15 Mouse_2a	781	GFPEALIPVQLLWVNLVTDGLPATALGFNPPDLDIMNKPPRNPKEPLISGWLFFRYLAIG
Mouse_2b	781
Rat_2b	781
Rat_2a	781
Dog_2a	781
20 Cat_2a	781
Pig_2a	781
Pig_2b	781
Human_2b	781
Human_2c	781
25 Human_2a	781
Rabbit_2a	781
Rabbit_2b	781
30 Mouse_2a	841	CYVGAATVGAAGWWFIAADGGPRVSFYQLSHFLQCKEDNPFDGVDCAIFESPYPMTMAL
Mouse_2b	841
Rat_2b	841
Rat_2a	841
Dog_2a	841
Cat_2a	841
35 Pig_2a	841
Pig_2b	841
Human_2b	841
Human_2c	841
Human_2a	841
40 Rabbit_2a	841
Rabbit_2b	841
45 Mouse_2a	901	SVLVTIEMCNALNSLSENQSLLRMPPWENIWLVGSICLMSMSLHFLILYVEPLPLIFQITP
Mouse_2b	901
Rat_2b	901
Rat_2a	901
Dog_2a	901
Cat_2a	901
Pig_2a	901
50 Pig_2b	901
Human_2b	901
Human_2c	901
Human_2a	901
Rabbit_2a	901
55 Rabbit_2b	901

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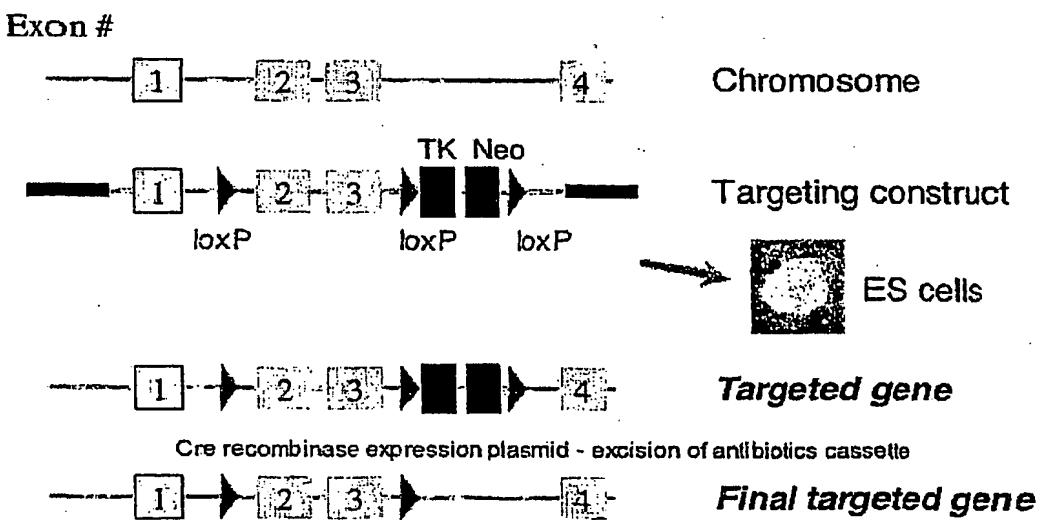
Mouse_2a	961	LNLTQWLMVLKISLPVILMDETLKFVARNYLEQPAILE-----
Mouse_2b	961GKECVQPATKSSCSLSACTDGISWPF
Rat_2b	961GKECA.....P.....
Rat_2a	961AILE
5 Dog_2a	961AILE
Cat_2a	961AILE
Pig_2a	961AILE
Pig_2b	961GKEC.....F.....
Human_2b	961	..V.....GKEC.....F.....
10 Human_2c	961	..V.....VLSSL
Human_2a	961	..V.....AILE
Rabbit_2a	961	..V.....AILE
Rabbit_2b	961	..V.....GKEC....PQ....W....E.V....
15 Mouse_2b	1021	VLLIMPLVVWVYSTDTNFSDMFWS
Rat_2b	1020
Pig_2b	1019
Human_2b	1019
Rabbit_2b	1019V...M.....LL..

Fig. 3 Targeting construct for Serca2 flox gene modification. Sequence information.

----- Serca2 gene -----
5 **LoxP site 1: Intron 1**
underlined sequence = loxP site and cloning sequence
Exon I (partial)
10 CACGGGGCTGAGCTTGGAGCAGGTCAAGAAGCTCAAGGGAGAGATGGGGCTCCAACGgtaggtgcggggcg
ccggggctgcagcgccgcggcgccggcccgagcgccaaggaaagatggctgaccggctccacctcgctggg
gcttggctcgccgcggcgcccgacggctgcgagagggccggcggtccacgcgcgggtctggccatcgccg
15 accttaggggtctcgatcaagcttacccgtcgatcgacccgtcgagggggggcccgatccgg
gataaattcgagctcgccgggatcgatccggaaaccctaataATAACTTCGTATAATGTATGCTATACG
AAGTTTATTtaggtccctcgacctcgagccaaagctccGGGGAtctcgagccgggtaccccccggccggcg
ctcagcgagtcggattggggggggggagagggagtggaaaggaggagggttctcgccgtggctg
agtccccccggatttatgaggcgctcgatgttggatggaaaccctcgaccgttctgtgtccccc
15 gttgcacatctggcagaagtgtatgaccctcgactgatgactgcattgttggaggccggagagggtt
cgggcagttccgaggccactgattaccagggtcgatccggggatccggaccgttctgtgtccccc
gttgcgttccatcacccatccggccattcagacaacgatgggtgtgaatttagcagtttataaaa
agcgctaatacataatcttcatttttttc
----- Serca2 gene -----
20 **LoxP site 2: Intron 3 5' of genomic XcmI site**
underlined sequence = loxPsite, cloning sites and partial HSV-TK
ccaattttattcttagaacattgtattcttatactgtgtatggaaatgtaaaaatcatacacttgc
ttagtttccaaaactgataactgtatggttcaattatgtattcacatcgatccgggg
25 GATCCggaaaccttaatATAACTTCGTATAATGTATGCTATACGAAAGTTTATTtaggtccctcgacccgtcg
cccaagctgatccctctcgatcgagcccaagctgtttccgcctcagaagccatagagccaccgc
cccaagcatgcctgtatgttcccaatctccccccttgcgttgcgcggccaccggccagaata
gaatgacacactactcgacaaatgcgtatgcatttcctatttatttagaaaggacagtggagtt
cttcaggggtcaaaaggcagccggggggggaaacaacacatgggtggcaactagaaggcacactcg
ggctgtatcgacgtctcgatcgatggaaattgtatccctcagaagaactcgatcgatggcgatagaaggc
30 gatgcgtgcgaatcgggagccggcgat*ccgtaaaggcagccgggg*ccggccattcgccqccaaac
tcttcagcaatatcacgggtagccacgtatgt*ctgataagcggtccggccacacccaa*ccggcca
caagtc*atgaaatcca*aaaaaggccggccatccggccacc*atgattt*ccggcaaggccctt*c
catgggtcaccgac*aga*catt*tccgt*c*ggcattgcgc*ccct
----- HSV-TK Neo antibiotics cassette -----
35 **LoxP site 3: Intron 3 5' of genomic XcmI site**
underlined sequence = loxP site, cloning sites and partial Neo gene
gttttcat*accaccqcgqggtcccggcc*gatat*tccacccgtc*ag*ccgtgtgtgtgggtgtaaatg
ttcgcgattgttgcqaaqcccc*agcaccggccagtaagtcatcggtcggtacgttagacgatatcg
40 cgccgcgaaaccacggggccaccacqcaagttcgatgggtgtgtttcccatcc*gtggggac*gtctatataa
acc*qcaagtgcgtggcatttctgcgtccggcgacttccgtggcttctgcgtccggcgaggccgaa
cgccgtacgtcggttgcataggccgcgagaacgcgcagccgtgcgaaacgcgcgggtgtatggccgg
ggtacgaaaggccatcgcgcttctacaaggcgctggccgaaagggtgcggggagtttacgcaccaagatct
45 gcggcacgctgtgacqctgtttaagcggtcgatcgatccgggtcgatcgatccggccacacgcgtcacc
ttaatatgcgaaatggacatcgacccgcgcggccggactcgatctgcgtgtcgatccggccatgcgatg
gacgctggccgggtttgtcgacatgggtggaaacatccaggccctgggtggagaggctttgtctcc
tcttgcaaaaccacactgctcgacattgggtggaaacatccaggccctgggtggagaggctttgtctcc
50 tcttgaaaaccacactqctcgatccggccaccctaataATAACTTCGTATAATGTATGCTATACGAA
GTAATtaggtccctcgacccgtcgagccaaagctgatccctcgatcgatccggccatggccatcgaaa
acatcgatggagg

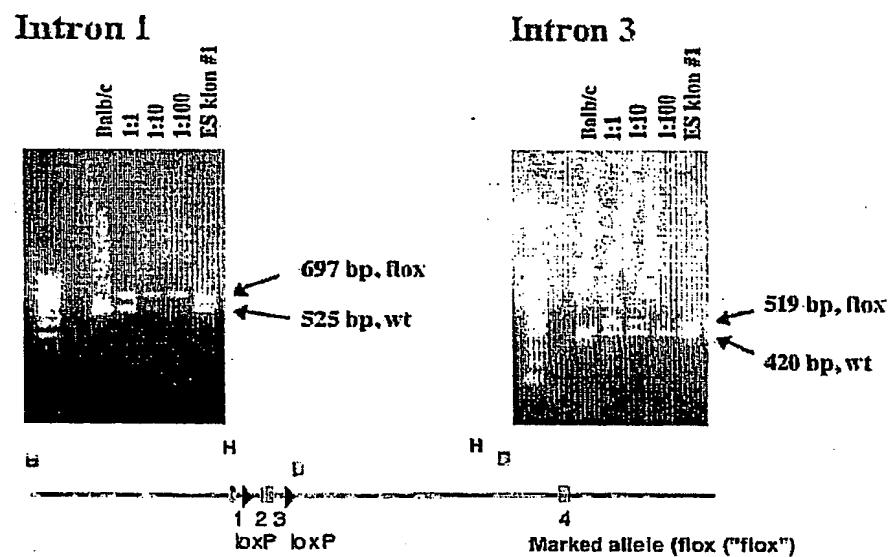
Fig. 4A Schematic representation of genetic manipulation.

Serca2 (*atp2a2*) gene modification



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Fig 4 B: Verification of Serca locus targeting events offspring from chimeric mice.



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Fig. 6 Cardiac ANP mRNA expression.

5

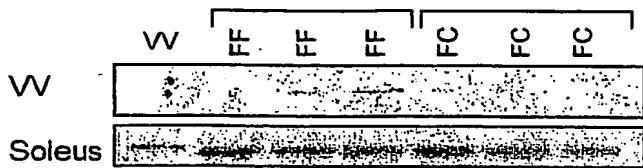


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Fig. 7 Serca2 protein expression.

5

A



B

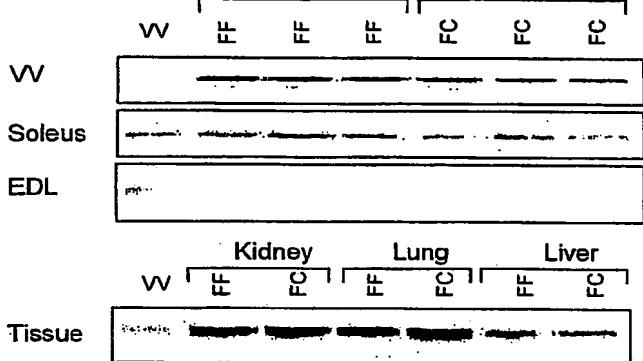
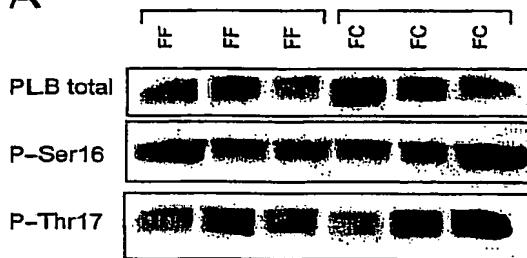
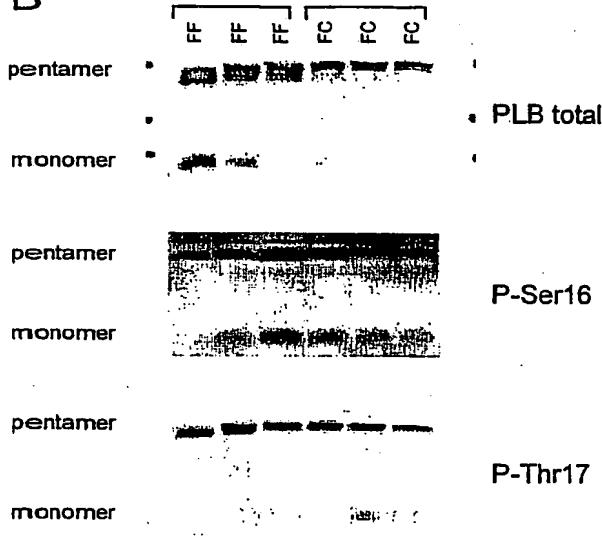


Fig. 8 Compensatory mechanisms in *Serca*^{flx} MLC-2v-Cre mice.

A



B



C

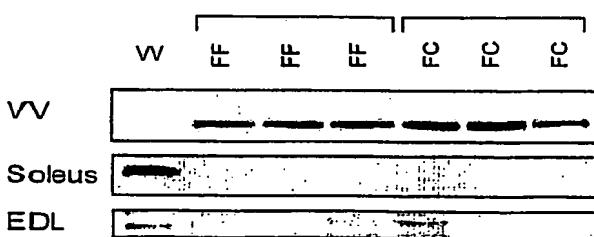
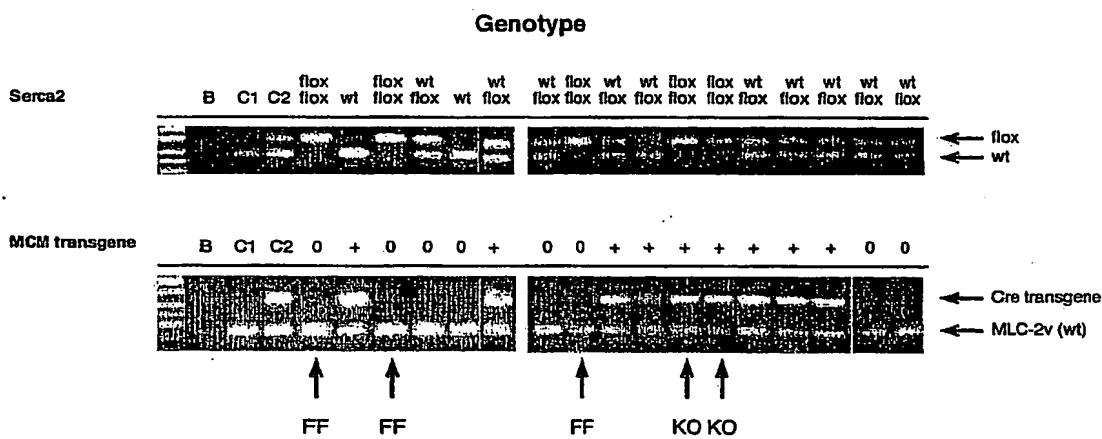


Figure 9 Genotypes PCR

5



10 Generation of animals with $\text{Serca2}^{\text{flox}}$ and MCM transgene alleles.

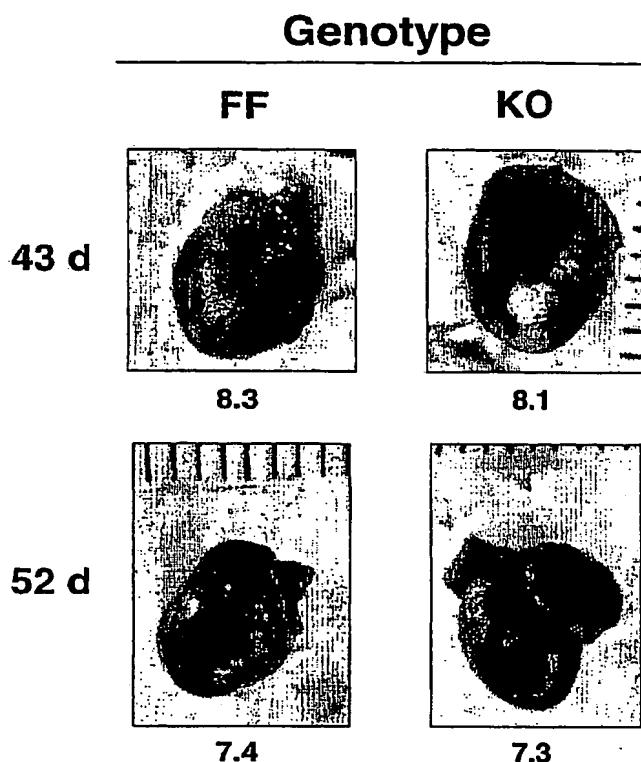
Genotypes FF, $\text{Serca2}^{\text{flox/flox}}$, KO, $\text{Serca2}^{\text{flox/flox}}$ MCM

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Figure 10

Heart morphology

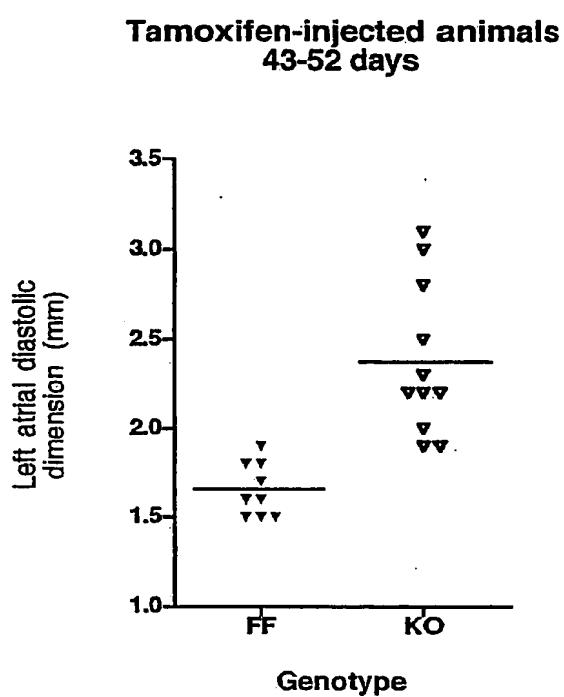
5



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Figure 11 Pilot series left atrial diastolic diameter.

5



10

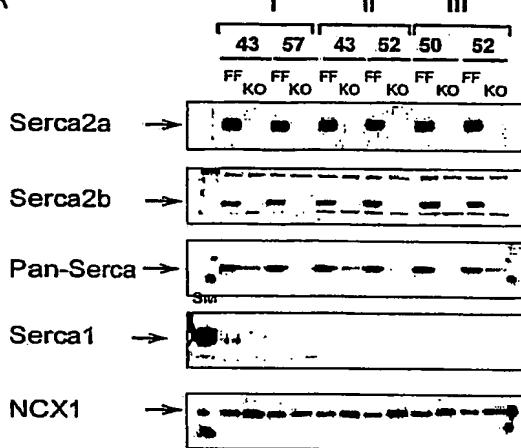
15

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Figure 12 Serca protein content in tamoxifen-induced FF and KO mice

5

A



B



10

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